



SEQUENCE LISTING

<110> Gijzen, Mark

<120> Soybean Seed Coat Peroxidase Structural Gene And Regulatory Region

<130> 76-105

<140> US 08/939,905

<141> 1996-09-30

<150> US 08/723,414

<151> 1996-09-30

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<170> PatentIn version 3.0

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Tyr Arg Glu Thr Cys Pro Asn Leu Phe Pro Ile Val Phe Gly Val Ile
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ttc gat gct tct ttc acc gat ccc cga atc ggg gcc agt ctc atg agg 192
Phe Asp Ala Ser Phe Thr Asp Pro Arg Ile Gly Ala Ser Leu Met Arg
50 55 60

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| ctg aac aac act gat aca ata gaa agc gag caa gat gca ctt cca aat Leu Asn Asn Thr Asp Thr Ile Glu Ser Glu Gln Asp Ala Leu Pro Asn 85 90 95 | 288 |
| atc aac tca ata aga gga ttg gac gtc aat gac atc aag aca gcg Ile Asn Ser Ile Arg Gly Leu Asp Val Val Asn Asp Ile Lys Thr Ala 100 105 110 | 336 |
| gtg gaa aat agt tgt cca gac aca gtt tct tgt gct gat att ctt gct Val Glu Asn Ser Cys Pro Asp Thr Val Ser Cys Ala Asp Ile Leu Ala 115 120 125 | 384 |
| att gca gct gaa ata gct tct gtt ctg gga gga ggt cca gga tgg cca Ile Ala Ala Glu Ile Ala Ser Val Leu Gly Gly Pro Gly Trp Pro 130 135 140 | 432 |
| gtt cca tta gga aga agg gac agc tta aca gca aac cya acc ctt gca Val Pro Leu Gly Arg Arg Asp Ser Leu Thr Ala Asn Arg Thr Leu Ala 145 150 155 160 | 480 |
| aatcaa aac ctt cca gca cct ttc aac ctc act caa ctt aaa gct Asn Gln Asn Leu Pro Ala Pro Phe Phe Asn Leu Thr Gln Leu Lys Ala 165 170 175 | 528 |
| tcc ttt gct gtt caa ggt ctc aac acc ctt gat tta gtt aca ctc tca Ser Phe Ala Val Gln Gly Leu Asn Thr Leu Asp Leu Val Thr Leu Ser 180 185 190 | 576 |
| ggt ggt cat acg ttt gga aga gct cgg tgc agt aca ttc ata aac cga Gly Gly His Thr Phe Gly Arg Ala Arg Cys Ser Thr Phe Ile Asn Arg 195 200 205 | 624 |
| tta tac aac ttc agc aac act gga aac cct gat cca act ctg aac aca Leu Tyr Asn Phe Ser Asn Thr Gly Asn Pro Asp Pro Thr Leu Asn Thr 210 215 220 | 672 |
| aca tac tta gaa gta ttg cgt gca aga tgc ccc cag aat gca act ggg Thr Tyr Leu Glu Val Leu Arg Ala Arg Cys Pro Gln Asn Ala Thr Gly 225 230 235 240 | 720 |
| gat aac ctc acc aat ttg gac ctg agc aca cct gat caa ttt gac aac Asp Asn Leu Thr Asn Leu Asp Leu Ser Thr Pro Asp Gln Phe Asp Asn 245 250 255 | 768 |
| aga tac tac tcc aat ctt ctg cag ctc aat ggc tta ctt cag agt gac Arg Tyr Tyr Ser Asn Leu Leu Gln Leu Asn Gly Leu Leu Gln Ser Asp 260 265 270 | 816 |
| caa gaa ctt ttc tcc act cct ggt gct gat acc att ccc att gtc aat Gln Glu Leu Phe Ser Thr Pro Gly Ala Asp Thr Ile Pro Ile Val Asn 275 280 285 | 864 |

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| agc ttc agc agt aac cag aat act ttc ttt tcc aac ttt aga gtt tca Ser Phe Ser Ser Asn Gln Asn Thr Phe Phe Ser Asn Phe Arg Val Ser 290 295 300 | 912 |
| atg ata aaa atg ggt aat att gga gtg ctg act ggg gat gaa gga gaa Met Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Asp Glu Gly Glu 305 310 315 320 | 960 |
| att cgc ttg caa tgt aat ttt gtg aat gga gac tcg ttt gga tta gct Ile Arg Leu Gln Cys Asn Phe Val Asn Gly Asp Ser Phe Gly Leu Ala 325 330 335 | 1008 |
| agt gtg gcg tcc aaa gat gct aaa caa aag ctt gtt gct caa tct aaa Ser Val Ala Ser Lys Asp Ala Lys Gln Lys Leu Val Ala Gln Ser Lys 340 345 350 | 1056 |
| taaaccaata attaatgggg atgtgcatac tagcttagcat gtaaaaggcaa attaggttgt aaacctcttt gctagctata ttgaaaataaaa ccaaaggagt agtgtgcatac tcaattcgat tttccatgt accttttggaa atattatgtt ataatttattt gaatctcttt aaggtaactta attaatca | 1116 1176 1236 1244 |

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| catatattag ctaaattagt tgttctaatt ggctatataa accctattgt actctttgta | 360 |
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| ctaatttgga gaatttgaat tatgatcatt aaatactcct ctcctgacta cttcgtccc | 660 |
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| tttaaaaaagt catacatgca aataatttt taatagttt cagttaaatt ttacagtaa | 840 |
| aaatgcataa aaattaaact ttatTTTCC aagtcatcat ttagtcaaatt cccaaaacaa | 900 |
| tgattatttt ttgcaaatga atgtttattg aacattttaa ttagtcaaatt ttaattctgg | 960 |
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| aaccatatta acgaagattt taattggat tcaagttcat gaacttagta aataagttt | 1140 |
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| gta gtg gca ttg ttg tgt gca ttt gct atg cat gca ggt ttt tca gtc | 1601 |
| Val Val Ala Leu Leu Cys Ala Phe Ala Met His Ala Gly Phe Ser Val | |
| 10 15 20 | |
| tct tat gct cag ctt act cct acg ttc tac aga gaa aca ttt cca aat | 1649 |
| Ser Tyr Ala Gln Leu Thr Pro Thr Phe Tyr Arg Glu Thr Cys Pro Asn | |
| 25 30 35 | |
| ctg ttc cct att gtg ttt gga gta atc ttc gat gct tct ttc acc gat | 1697 |
| Leu Phe Pro Ile Val Phe Gly Val Ile Phe Asp Ala Ser Phe Thr Asp | |
| 40 45 50 55 | |
| ccc cga atc ggg gcc agt ctc atg agg ctt cat ttt cat gat tgc ttt | 1745 |
| Pro Arg Ile Gly Ala Ser Leu Met Arg Leu His Phe His Asp Cys Phe | |
| 60 65 70 | |
| gtt caa gtacgtactt tttttttcc ttccaaaatg ccctgcataat ttaacaagat | 1801 |
| Val Gln | |
| tgctttgttc acctagaaaa atgtgtttt ttcaacgatc ttacgtacgt ttgtttgggt | 1861 |
| tgaaaaataa atcagaaaa gatcaagaaa atagctagaa agaaagcaac gtttttttaa | 1921 |
| aaggtattta gtgtgagaaa aatattaaaa ctgaagagaa agaaattaaa taagctttc | 1981 |
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| tgatctgaac aaattaagtt gttatattt cattgtgaca g ggt tgt gat gga tca Gly Cys Asp Gly Ser | 2397 |
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| gtt ttg ctg aac aac act gat aca ata gaa agc gag caa gat gca ctt Val Leu Leu Asn Asn Thr Asp Thr Ile Glu Ser Glu Gln Asp Ala Leu | 2415 |
| 80 85 90 | |
| cca aat atc aac tca ata aga gga ttg gac gtt gtc aat gac atc aag Pro Asn Ile Asn Ser Ile Arg Gly Leu Asp Val Val Asn Asp Ile Lys | 2493 |
| 95 100 105 110 | |
| aca gcg gtg gaa aat agt tgt cca gac aca gtt tct tgt gct gat att Thr Ala Val Glu Asn Ser Cys Pro Asp Thr Val Ser Cys Ala Asp Ile | 2541 |
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| aac cga acc ctt gca aatcaa aac ctt cca gca cct ttc ttc aac ctc Asn Arg Thr Leu Ala Asn Gln Asn Leu Pro Ala Pro Phe Phe Asn Leu 160 165 170 | 3706 |
| act caa ctt aaa gct tcc ttt gct gtt caa ggt ctc aac acc ctt gat Thr Gln Leu Lys Ala Ser Phe Ala Val Gln Gly Leu Asn Thr Leu Asp 175 180 185 | 3754 |
| tta gtt aca ctc tca ggtatacata atcaattttt tatttgctat tagctagcaa Leu Val Thr Leu Ser 190 | 3809 |
| taaaaaagtct ctgatacaga catattnaga taaattaatt tctccataaa catttataat aaaattatca atttatgtac ttaaaaattt tggattgaag ctctttcat ccaactttta ctaaagttaa ggtgcataataataaaaata aactatctct tgtttcttat aaaaagattg aagataagtt aaagtctact tataaatcat taatatatgt ata ggt ggt cat acg Gly Gly His Thr 195 | 3869 3929 3989 4044 |
| ttt gga aga gct cgg tgc agt aca ttc ata aac cga tta tac aac ttc Phe Gly Arg Ala Arg Cys Ser Thr Phe Ile Asn Arg Leu Tyr Asn Phe 200 205 210 | 4092 |
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| 325 330 335 340 | |
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| 345 350 | |
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gacaaaagga gagattagaa aacaatgcaa ctttgtgaac tttgtgaact caaattctgc 1020
agaacttagat ttagccacca tagcatccat agtagaatca ttagaggatg gtattgctag 1080
tgtaatataa ataaatttagc gtaaaatgcac ttattgaaat cttgtgacta gatgccacta 1140
ataaataagt tataactagg cacatttcat gtcacttgaa atttcatgcc tgttatatgag 1200

<210> 13

<211> 1200

<212> DNA

<213> *Medicago sativa*

<400> 13
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ttcctcagat gcacaactta gtcccacttt ttacagcaaa acgtgtccaa ctgttagttc 120

cattgttagc aatgtcttaa caaacgtttc taagacagat cctcgcatgc ttgctagtct 180
cgtcaggc tt cactttcatg actgtttgt tctggatgt gatgcctcag ttttgctgaa 240
caatactgct acaatcgtaa gcgacaaca a agctttcca aataacaact ctctaagggg 300
tttggatgtt gtgaatcaga tcaaaaactgc tgttagaaagt gcttgccta acacagttc 360
tttgtgctgat attcttcac ttgctcaagc atcctctgtt ctggcacaag gtcctagttg 420
gacggttcct ttaggaagaa gggatggttt aaccgcaa ac cgaacacttg caaatcaaaa 480
tcttccggct ccattcaatt ctttggatca cttttaactg catttgactg ctcaaggcct 540
cattactcct gttcttagttg ccctctcggg tgctcataca tttggaagag ctcattgcgc 600
acaatttgtt agtcgattgt acaacttcag cagtactgga agtcccgtc caactcttaa 660
cacaacttac ttacaacaac tgccgacaat atgtcccaat ggtggacctg gcacaaacct 720
taccaatttc gatccaacga ctccgtataa atttgacaag aactattact ccaatcttca 780
agtaaaaaag gtttgctcc aaagtgtatca agagttgtc tcaacttctg gtgcagatac 840
cattagcatt gtcgacaaat tcagcaccga tcaaaaatgct ttcttgaga gctttaaggc 900
tgcaatgatt aaaatggca atattgggt gctaacaggg acaaaaggag agattagaaa 960
acaatgcaac tttgtgaact caaattctgc agaactagat ttagccacca tagcatccat 1020
agtagaatca ttagaggatg gaattgctag tgtaatataa ataaatttagc gaaaatgcac 1080
ttattgaaat cttgtgacta gatcccacta ataaataagt tataactagg cacattcat 1140
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<210> 14

<211> 1200

<212> DNA

<213> *Medicago sativa*

<400> 14
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accctttca aatgcacaac tagatccttc attttacaac agtacatgtt ctaatcttga 120
ttcaatcgta cgtgggtgc tcacaaatgt ttcacaatct gatcccagaa tgcttggtag 180
tctcatcagg ctacatttc atgactgtt tttcaaggt tgccatgcct cgattttgtc 240
gaacgatacg gctacaatag tgagcgagca aagtgcacca ccaaataaca actccataag 300

| | |
|--|------|
| aggttggat gtgataaacc agatcaaaac agcggtggaa aatgcttgc ctaacacagt | 360 |
| ttcttgtct gatattcttgc ctcttctgc taaaatatca tctgatctgg caaatggtcc | 420 |
| tacttggcaa gttccattag gaagaaggga tagttgaca gcaaataatt cccttgagc | 480 |
| tcaaaatctt cctgccccca ctttcaacct tactcgacta aaatctaact ttgataatca | 540 |
| aaacacctgact actactgatc tagttgact ctcaggtggc catacaattg gaagaggtca | 600 |
| atgcagatt ttcgttgatc gattatacaa tttcagcaac actggaaacc ccgattcaac | 660 |
| tcttaacacg acctatttac aaacattgca agcaatatgt cccaatggtg gacctggtagc | 720 |
| aaacctaacc gatttggacc caaccacacc agatacattt gactccaact actactccaa | 780 |
| tctccaagtt ggaaagggtt tgttcagag tgaccaagag ctttttcca gaaatggttc | 840 |
| tgacactatt tctattgtca atagttcgc caataatcaa actctcttct ttgaaaattt | 900 |
| tgtacactt atgataaaaa tggtaataat tggagttta actggatctc aaggtgaaat | 960 |
| tagaacacag tgtaatgctg tgaatggaa ttcttctgga ttggctactg tagtcaccaa | 1020 |
| agaatcatca gaagatggaa tggctagctc attctaaata taagcttggaa aaatattgaa | 1080 |
| gaggttctat aattttgtgc atacatataat ggtatgtca tgtggtgtat tatgttttg | 1140 |
| ttatgttctt caagttgatc agggactgta gaagctccct aataatattt gtgtcaaagt | 1200 |

<210> 15

<211> 283

<212> PRT

<213> Glycine max

<400> 15

| | | | |
|---|---|----|----|
| Phe His Asp Cys Phe Val Gln Gly Cys Asp Gly Ser Val Leu Leu Asn | | | |
| 1 | 5 | 10 | 15 |

| | | | |
|---|----|----|--|
| Asn Thr Asp Thr Ile Glu Ser Glu Gln Asp Ala Leu Pro Asn Ile Asn | | | |
| 20 | 25 | 30 | |

| | | | |
|---|----|----|--|
| Ser Ile Arg Gly Leu Asp Val Val Asn Asp Ile Lys Thr Ala Val Glu | | | |
| 35 | 40 | 45 | |

| | | | |
|---|----|----|--|
| Asn Ser Cys Pro Asp Thr Val Ser Cys Ala Asp Ile Leu Ala Ile Ala | | | |
| 50 | 55 | 60 | |

| | |
|---|--|
| Ala Glu Ile Ala Ser Val Ala Gly Arg Arg Ser Gly Trp Pro Val Pro | |
|---|--|

65 70 75 80
 Leu Gly Arg Arg Asp Ser Leu Thr Ala Asn Arg Thr Leu Ala Asn Gln
 85 90 95
 Asn Leu Pro Ala Pro Phe Phe Asn Leu Thr Gln Leu Lys Ala Ser Phe
 100 105 110
 Ala Val Gln Gly Leu Asn Thr Leu Asp Leu Val Thr Leu Ser Gly Gly
 115 120 125
 His Thr Ser Gly Arg Ala Arg Cys Ser Thr Phe Ile Asn Arg Leu Tyr
 130 135 140
 Asn Phe Ser Asn Thr Gly Leu Ile His Leu Asp Thr Thr Tyr Leu Glu
 145 150 155 160
 Val Leu Arg Ala Arg Cys Pro Gln Asn Ala Thr Gly Asp Asn Leu Thr
 165 170 175
 Asr. Leu Asp Leu Ser Thr Pro Asp Gln Phe Asp Asn Arg Tyr Tyr Ser
 180 185 190
 Asr. Leu Leu Gln Leu Asn Gly Leu Leu Gln Ser Asp Gln Glu Arg Phe
 195 200 205
 Ser Thr Pro Gly Ala Asp Thr Ile Pro Leu Ser Ile Ala Ser Ala Asn
 210 215 220
 Gln Asn Thr Phe Phe Ser Asn Phe Arg Val Ser Met Ile Lys Met Gly
 225 230 235 240
 Asn Ile Gly Val Leu Thr Gly Asp Glu Gly Glu Ile Arg Leu Gln Cys
 245 250 255
 Asn Phe Val Asn Gly Asp Ser Phe Gly Leu Ala Ser Val Ala Ser Lys
 260 265 270
 Asp Ala Lys Gln Lys Leu Val Ala Gln Ser Lys
 275 280
 <210> 16
 <211> 355
 <212> PRT
 <213> *Medicago sativa*
 <400> 16
 Met Asn Ser Leu Arg Ala Val Ala Ile Ala Leu Cys Cys Ile Val Val
 1 5 10 15
 Val Leu Gly Gly Leu Pro Phe Ser Ser Asn Ala Gln Leu Asp Pro Ser

325

330

335

Gly Leu Ile Asn Val Ala Ser Ala Asp Ser Ser Glu Glu Gly Met Val
340 345 350

Ser Ser Met
355

<210> 17

<211> 358

<212> PRT

<213> *Medicago sativa*

<400> 17

1
 Leu Gly Gly Leu Pro Phe Ser Ser Asp Ala Gln Leu Ser Pro Thr Phe
 20 25 30

Tyr Ser Lys Thr Cys Pro Thr Val Ser Ser Ile Val Ser Asn Val Leu
 35 40 45

59
Thr Asn Val Ser Lys Thr Asp Pro Arg Met Leu Ala Ser Leu Val Arg
50 55 60

Asn Asn Ser Leu Arg Gly Leu Asp Val Val Asn Gln Ile Lys Leu Ala
 100 105 110

100 Val Glu Val Pro Cys Pro Asn Thr Val Ser Cys Ala Asp Ile Leu Ala
105 120 125

115 120
 Leu Ala Ala Gln Ala Ser Ser Val Leu Ala Gln Gly Pro Ser Trp Thr
 135 140

130 135
Val Pro Leu Gly Arg Arg Asp Gly Leu Thr Ala Asn Arg Thr Leu Ala
150 155 160

145 150
Asn Gln Asn Leu Pro Ala Pro Phe Asn Ser Leu Asp Gln Leu Lys Ala
 155 170 175

165 170
Ala Phe Thr Ala Gln Gly Leu Asn Thr Thr Asp Leu Val Ala Leu Ser
 185 190

180 185
... Ala His Thr Phe Gly Arg Ala His Cys Ala Gln Phe Val Ser Arg

195 200 205
Leu Tyr Asn Phe Ser Ser Thr Gly Ser Pro Asp Pro Thr Leu Asn Thr
210 215 220
Thr Tyr Leu Gln Gln Leu Arg Thr Ile Cys Pro Asn Gly Gly Pro Gly
225 230 235 240
Thr Asn Leu Thr Asn Phe Asp Pro Thr Thr Pro Asp Lys Phe Asp Lys
245 250 255
Asn Tyr Tyr Ser Asn Leu Gln Val Lys Lys Gly Leu Leu Gln Ser Asp
260 265 270
Gln Glu Leu Phe Ser Thr Ser Gly Ala Asp Thr Ile Ser Ile Val Asn
275 280 285
Lys Phe Ser Thr Asp Gln Asn Ala Phe Phe Glu Ser Phe Lys Ala Ala
290 295 300
Met Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Thr Lys Gly Glu
305 310 315 320
Ile Arg Lys Gln Cys Asn Phe Val Asn Phe Val Asn Ser Asn Ser Ala
325 330 335
Glu Leu Asp Leu Ala Thr Ile Ala Ser Ile Val Glu Ser Leu Glu Asp
340 345 350
Gly Ile Ala Ser Val Ile
355

<210> 18

<211> 347

<212> PRT

<213> *Medicago sativa*

<400> 18

Met Trp Cys Val Val Leu Leu Val Val Leu Gly Gly Leu Pro Phe Ser
1 5 10 15
Ser Asp Ala Gln Leu Ser Pro Thr Phe Tyr Ser Lys Thr Cys Pro Thr

20 25 30
Val Ser Ser Ile Val Ser Asn Val Leu Thr Asn Val Ser Lys Thr Asp

35 40 45
Pro Arg Met Leu Ala Ser Leu Val Arg Leu His Phe His Asp Cys Phe
50 55 60

Val Leu Gly Cys Asp Ala Ser Val Leu Leu Asn Asn Thr Ala Thr Ile

| | | | |
|--|-----|----|-----|
| 65 | 70 | 75 | 80 |
| Val Ser Glu Gln Gln Ala Phe Pro Asn Asn Asn Ser Leu Arg Gly Leu | | | |
| 85 | 90 | | 95 |
| Asp Val Val Asn Gln Ile Lys Thr Ala Val Glu Ser Ala Cys Pro Asn | | | |
| 100 | 105 | | 110 |
| Thr Val Ser Cys Ala Asp Ile Leu Ala Leu Ala Gln Ala Ser Ser Val | | | |
| 115 | 120 | | 125 |
| Leu Ala Gin Gly Pro Ser Trip Thr Val Pro Leu Gly Arg Arg Asp Gly | | | |
| 130 | 135 | | 140 |
| Leu Thr Ala Asn Arg Thr Leu Ala Asn Gln Asn Leu Pro Ala Pro Phe | | | |
| 145 | 150 | | 160 |
| Asn Ser Leu Asp His Leu Lys Leu His Leu Thr Ala Gln Gly Leu Ile | | | |
| 165 | 170 | | 175 |
| Thr Pro Val Leu Val Ala Leu Ser Gly Ala His Thr Phe Gly Arg Ala | | | |
| 180 | 185 | | 190 |
| His Cys Ala Gln Phe Val Ser Arg Leu Tyr Asn Phe Ser Ser Thr Gly | | | |
| 195 | 200 | | 205 |
| Ser Pro Asp Pro Thr Leu Asn Thr Thr Tyr Leu Gln Gln Leu Arg Thr | | | |
| 210 | 215 | | 220 |
| Ile Cys Pro Asn Gly Gly Pro Gly Thr Asn Leu Thr Asn Phe Asp Pro | | | |
| 225 | 230 | | 240 |
| Thr Thr Pro Asp Lys Phe Asp Lys Asn Tyr Tyr Ser Asn Leu Gln Val | | | |
| 245 | 250 | | 255 |
| Lys Lys Gly Leu Leu Gln Ser Asp Gln Glu Leu Phe Ser Thr Ser Gly | | | |
| 260 | 265 | | 270 |
| Ala Asp Thr Ile Ser Ile Val Asp Lys Phe Ser Thr Asp Gln Asn Ala | | | |
| 275 | 280 | | 285 |
| Phe Phe Glu Ser Phe Lys Ala Ala Met Ile Lys Met Gly Asn Ile Gly | | | |
| 290 | 295 | | 300 |
| Val Leu Thr Gly Thr Lys Gly Glu Ile Arg Lys Gln Cys Asn Phe Val | | | |
| 305 | 310 | | 320 |
| Asn Ser Asn Ser Ala Glu Leu Asp Leu Ala Thr Ile Ala Ser Ile Val | | | |
| 325 | 330 | | 335 |
| Glu Ser Leu Glu Asp Gly Ile Ala Ser Val Ile | | | |
| 340 | 345 | | |

<210> 19

<211> 351

<212> PRT

<213> Medicago sativa

<400> 19

Met Leu Gly Leu Ser Ala Thr Ala Phe Cys Cys Met Val Phe Val Leu
1 5 10 15

Ile Gly Gly Val Pro Phe Ser Asn Ala Gln Leu Asp Pro Ser Phe Tyr
20 25 30

Asn Ser Thr Cys Ser Asn Leu Asp Ser Ile Val Arg Gly Val Leu Thr
35 40 45

Asn Val Ser Gln Ser Asp Pro Arg Met Leu Gly Ser Leu Ile Arg Leu
50 55 60

His Phe His Asp Cys Phe Val Gln Gly Cys Asp Ala Ser Ile Leu Leu
65 70 75 80

Asn Asp Thr Ala Thr Ile Val Ser Glu Gln Ser Ala Pro Pro Asn Asn
85 90 95

Asn Ser Ile Arg Gly Leu Asp Val Ile Asn Gln Ile Lys Thr Ala Val
100 105 110

Glu Asn Ala Cys Pro Asn Thr Val Ser Cys Ala Asp Ile Leu Ala Leu
115 120 125

Ser Ala Glu Ile Ser Ser Asp Leu Ala Asn Gly Pro Thr Trp Gln Val
130 135 140

Pro Leu Gly Arg Arg Asp Ser Leu Thr Ala Asn Asn Ser Leu Ala Ala
145 150 155 160

Gln Asn Leu Pro Ala Pro Thr Phe Asn Leu Thr Arg Leu Lys Ser Asn
165 170 175

Phe Asp Asn Gln Asn Leu Ser Thr Thr Asp Leu Val Ala Leu Ser Gly
180 185 190

Gly His Thr Ile Gly Arg Gly Gln Cys Arg Phe Phe Val Asp Arg Leu
195 200 205

Tyr Asn Phe Ser Asn Thr Gly Asn Pro Asp Ser Thr Leu Asn Thr Thr
210 215 220

Tyr Leu Gln Thr Leu Gln Ala Ile Cys Pro Asn Gly Gly Pro Gly Thr
225 230 235 240

Asn Leu Thr Asp Leu Asp Pro Thr Thr Pro Asp Thr Phe Asp Ser Asn
245 250 255

Tyr Tyr Ser Asn Leu Gln Val Gly Lys Gly Leu Phe Gln Ser Asp Gln

260

265

270

Glu Leu Phe Ser Arg Asn Gly Ser Asp Thr Ile Ser Ile Val Asn Ser
275 280 285

Phe Ala Asn Asn Gln Thr Leu Phe Phe Glu Asn Phe Val Ala Ser Met
290 295 300

Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Ser Gln Gly Glu Ile
305 310 315 320

Arg Thr Gln Cys Asn Ala Val Asn Gly Asn Ser Ser Gly Leu Ala Thr
325 330 335

Val Val Thr Lys Glu Ser Ser Glu Asp Gly Met Ala Ser Ser Phe
340 345 350

<210> 20

<211> 22

<212> DNA

<213> *Medicago sativa*

<400> 20

taaaatcata tcagcttact cc

22